# organic papers

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#### Key indicators

Single-crystal X-ray study T = 225 KMean  $\sigma$ (C–C) = 0.003 Å R factor = 0.057 wR factor = 0.173 Data-to-parameter ratio = 14.3

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# Tetrakis(2-methoxy-5-pyridyl)silane

The title compound,  $C_{24}H_{24}N_4O_4Si$ , crystallizes from propionic acid–hexane to produce a close-packed structure with no included guests. In contrast, crystallization of the closely related tetrakis(1,2-dihydro-2-oxo-5-pyridyl)silane under similar conditions is directed by hydrogen bonding of the pyridinone groups to yield an open diamondoid network, which is filled by a combination of interpenetration and inclusion of guests.

### Comment

A promising strategy in crystal engineering is based on the use of molecules that can associate by forming reliable directional interactions, thereby placing their neighbors in predictable positions. Such molecules, which have been called tectons, from the Greek word for builder (Simard et al., 1991), can serve as the basic elements of a rational nanoscale construction set for building crystalline networks with predetermined architectures and properties. The directional interactions that hold such structures together cannot normally be optimized when the tectons are closely packed, so the resulting networks are typically porous, with interiors filled by interpenetration of independent networks or by the inclusion of guests (Saied, Maris, Wang et al., 2005; Malek, Maris, Perron & Wuest, 2005; Malek, Maris, Simard & Wuest, 2005). For example, crystallization of tetrakis(1,2-dihydro-2-oxo-5-pyridyl)silane, (I), is directed by characteristic hydrogen bonding of its pyridinone groups to give open interpenetrated diamondoid networks, in which approximately 60% of the volume remains available for the inclusion of guests (Saied et al., 2003).



In contrast, the structure of the title compound, (II), is close-packed, devoid of guests, and held together primarily by a combination of van der Waals interactions, aromatic interactions,  $C-H\cdots O$  interactions, and  $C-H\cdots N$  interactions (Figs. 1 and 2). Two  $C-H\cdots N$  interactions [ $H\cdots N$  2.68 (1) Å] are formed with the closest neighbor, which is separated by an Si $\cdots$ Si distance of 6.395 (3) Å. The average C-Si-C angle (109.46°) is very close to the ideal tetrahedral value.

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As expected, the 2-methoxy-5-pyridyl groups of the title compound, (II), adopt a near-planar *syn* conformation, (III), in which the methoxy substituent lies close to the mean plane of the pyridine ring, and the methyl group is oriented toward the N atom. In this structure, the average N-C-O-C dihedral angle is 2.8°. Analysis of all 42 structures in the Cambridge Structural Database (Version 5.26 of February 2005; Allen, 2002) containing one or more 5-substituted 2-methoxy-5-pyridyl groups revealed only one example of the alternative planar *anti* conformation, (IV).

Together, our observations are a valuable addition to the growing body of evidence showing how the crystallization of tectons differs distinctively from that of close structural analogs lacking multiple sites of strong directional interaction (Saied, Maris, Simard & Wuest, 2005). Tecton (I) forms open hydrogen-bonded networks with substantial porosity, whereas its *O*-methyl derivative, (II), crystallizes under very similar conditions as a close-packed structure with no included guests. The divergent behavior of compounds (I) and (II) illustrates how suitably oriented hydrogen-bonding groups can have a profound effect as directors of crystallization.

## Experimental

Tetrakis(2-methoxy-5-pyridyl)silane, (II), was prepared from tetrakis(1,2-dihydro-2-oxo-5-pyridyl)silane, (I), by a general method for the *O*-alkylation of pyridinones (Comins & Jianhua, 1994). Crystals were grown by allowing hexane to diffuse slowly into a solution of the title compound, (II), in propionic acid.

#### Crystal data

C <sub>24</sub> H <sub>24</sub> N <sub>4</sub> O <sub>4</sub> Si
$M_r = 460.56$
Triclinic, P1
a = 9.661 (3)  Å
b = 10.364 (4) Å
c = 12.230 (6) Å
$\alpha = 107.74 \ (4)^{\circ}$
$\beta = 94.68 \ (3)^{\circ}$
$\gamma = 99.22 \ (3)^{\circ}$
$V = 1140.1 (9) \text{ Å}^3$

## Data collection

Enraf-Nonius CAD-4 diffractometer  $\omega/2\theta$  scans Absorption correction: analytical (ABSORP in *NRCVAX*; Gabe *et al.*, 1989)  $T_{\min} = 0.570, T_{\max} = 0.810$ 8614 measured reflections 4319 independent reflections

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.057$   $wR(F^2) = 0.173$  S = 1.114319 reflections 302 parameters H-atom parameters constrained Z = 2  $D_x = 1.342 \text{ Mg m}^{-3}$ Cu  $K\alpha$  radiation Cell parameters from 25 reflections  $\theta = 15.0-30.0^{\circ}$   $\mu = 1.24 \text{ mm}^{-1}$  T = 225 (2) K Block, colorless  $0.52 \times 0.48 \times 0.18 \text{ mm}$ 

3964 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.078$   $\theta_{max} = 69.8^{\circ}$   $h = -11 \rightarrow 11$   $k = -12 \rightarrow 12$   $l = -14 \rightarrow 14$ 6 standard reflections frequency: 60 min intensity decay: 0.1%

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.1135P)^2 \\ &+ 0.4013P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.43 \ {\rm e}\ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.53 \ {\rm e}\ {\rm \AA}^{-3} \\ {\rm Extinction \ correction: \ SHELXL97} \\ &({\rm Sheldrick, \ 1997}) \\ {\rm Extinction \ coefficient: \ 0.0133 \ (15)} \end{split}$$



#### Figure 1

A view of the structure of (II), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are represented by spheres of arbitrary radii.



Figure 2

Views of one molecule of (II) (red) with the neighbor (light blue) that is separated from it by an Si $\cdots$ Si distance of 6.395 (3) Å. The two molecules illustrated are related by a center of inversion and together constitute the contents of one unit cell. The left-hand view shows the closest contacts as broken lines, with an H $\cdots$ N distance of 2.68 (1) Å. In the right-hand view, the molecules are shown as CPK representations.

# Table 1 Selected geometric parameters (Å, °).

Si-C11	1.867 (2)	Si-C21	1.872 (2)
Si-C31	1.871 (2)	Si-C41	1.874 (2)
C11-Si-C31	110.28 (10)	C11-Si-C41	105.43 (10)
C11-Si-C21	110.38 (10)	C31-Si-C41	111.96 (10)
C31-Si-C21	109.15 (10)	C21-Si-C41	109.59 (10)
C17 - O1 - C14 - N13	81(3)	C37-O3-C34-N33	-0.6(3)
$C_{27} = O_{2} = C_{24} = N_{23}$	43(3)	C47 - O4 - C44 - N43	-0.7(3)
		0 0 011 IAB	017 (0)

After confirmation of their location in a difference map, H atoms were positioned geometrically and refined using a riding model, with C-H distances in the range 0.94–0.97 Å, and with  $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm C})$  for methyl groups or  $1.2U_{\rm eq}({\rm C})$  for other atoms.

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Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: modified versions of *NRC-2/NRC2A* (Ahmed *et al.*, 1973); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *enCIFer* (Allen *et al.*, 2004).

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